

## **STIC Biotechnology Systems Branch**

### **RAW SEQUENCE LISTING** **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/562,601  
Source: FWO  
Date Processed by STIC: 3/5/07

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

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<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

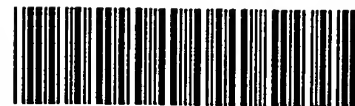
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Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

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1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):  
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06



IFWO

## RAW SEQUENCE LISTING

DATE: 03/05/2007

PATENT APPLICATION: US/10/562,601

TIME: 13:44:39

Input Set : N:\efs\03\_05\_07\10562601\_efs\sequencelisting.txt

Output Set: N:\CRF4\03052007\J562601.raw

3 <110> APPLICANT: Hengst den, Christiaan D.  
 4 Gajic, Olivera  
 5 Kuipers, Oscar P.  
 6 Kok, Jan  
 7 Sikkema, Jan  
 8 Geurts, Johannes M.W.  
 9 Nauta, Arjen  
 11 <120> TITLE OF INVENTION: Methods and means for regulating gene expression  
 13 <130> FILE REFERENCE: P63590US00  
 15 <140> CURRENT APPLICATION NUMBER: US 10/562,601  
 16 <141> CURRENT FILING DATE: 2005-12-28  
 18 <150> PRIOR APPLICATION NUMBER: PCT/NL2004/000474  
 19 <151> PRIOR FILING DATE: 2004-07-02  
 21 <150> PRIOR APPLICATION NUMBER: EP 03077074.7  
 22 <151> PRIOR FILING DATE: 2003-07-02  
 24 <160> NUMBER OF SEQ ID NOS: 234  
 27 <170> SOFTWARE: PatentIn Ver. 3.3

Does Not Comply  
Corrected Diskette Needed

see p. 2, 4

## ERRORED SEQUENCES

2299 <210> SEQ ID NO: 179  
 2300 <211> LENGTH: 26  
 2301 <212> TYPE: DNA  
 2302 <213> ORGANISM: Artificial Sequence  
 2304 <220> FEATURE:  
 2305 <223> OTHER INFORMATION: presence of the putative CodY box in arcA  
 E--> 2307 <400> SEQUENCE: 179 see p. 2  
 2308 atatttttttg aaaaattttt taaaaa

26

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<210> 179  
<211> 26  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> presence of the putative CodY box in arcA

Q<400> 179  
atatttttttg aaaaattttt taaaaa

26

delete extra opening bracket

## VERIFICATION SUMMARY

DATE: 03/05/2007

PATENT APPLICATION: US/10/562,601

TIME: 13:44:40

Input Set : N:\efs\03\_05\_07\10562601\_efs\sequencelisting.txt

Output Set: N:\CRF4\03052007\J562601.raw

L:2307 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:179 differs:178  
L:2739 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:214,Line#:2732  
L:2744 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:214,Line#:2737  
L:2750 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:214,Line#:2742  
L:2750 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:214,Line#:2747  
L:2914 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:227 after pos.:0  
L:2931 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:228 after pos.:0  
L:2960 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:230 after pos.:0

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<210> 15  
<211> 8  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> consensus sequence GTP binding motif in CodY homologs

<220>  
<221> Variant  
<222> (2)..(2)  
<223> Arg

if the amino acid at this location can represent another amino acid, use "Xaa" instead, and

specify in <2207-22237 section

<220>  
<221> Variant  
<222> (3)..(3) same  
<223> Met, Gly, Ile, Lys, Gln

<220>  
<221> Variant  
<222> (4)..(4) same  
<223> Gly

<220>  
<221> Variant  
<222> (5)..(5) same  
<223> Thr

<220>  
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<222> (7)..(7) same  
<223> Ser

<400> 15  
Gly Gly Glu Arg Leu Gly Thr Thr  
1 5